

M A L L A R I L R A G L R P A P E R G G SEQ ID NO:2  
 ATG GCC CTG CTG GCG CGG ATC CTG AGA GCC GGG CTG CGG CCG GCG CCC GAG CGG GGT GGG SEQ ID NO:1  
 ↑SEQ ID NO:3→  
 L L G G G A P R R P Q P A G A R L P A G 40  
 CTC CTG GGC GGC GGG GCC CCG CGG CGG CCT CAA CCC GCG GGC GCA CGG CTC CCG GCG GGG 120  
  
 A R A E D K G A G R P G S P P G G G R A 60  
 GCG CGG GCC GAG GAC AAA GGC GCC GGG CGG CCG GGG TCG CCG CCG GGA GGG GGC CGA GCC 180  
  
 E G P R S L A A M P G P R T L A N L A E 80  
 GAG GGT CCC CGG AGC CTC GCC GCC ATG CCG GGG CCG AGG ACC CTC GCC AAC CTG GCG GAG 240  
  
 F F C R D G F S R I H E I Q Q K H T R E 100  
 TTC TTC TGC AGG GAC GGC TTC AGC CGC ATC CAC GAG ATC CAG CAG AAG CAC ACA CGG GAA 300  
  
 Y G K I F K S H F G P Q F V V S I A D R 120  
 TAT GGA AAA ATC TTC AAG TCT CAC TTT GGT CCT CAG TTT GTA GTA TCT ATT GCA GAC CGC 360  
  
 D M V A Q V L R A E G A A P Q R A N M E 140  
 GAT ATG GTG GCT CAG GTG CTC CGG GCG GAG GGC GCT GCG CCC CAG AGA GCC AAC ATG GAG 420  
  
 S W R E Y R D L R G R A T G L I S A E G 160  
 TCC TGG CGG GAG TAC CGA GAC TTG CGG GGG AGA GCC ACC GGG CTC ATC TCG GCG GAG GGT 480  
  
 E Q W L K M R S V L R Q R I L K P K D V 180  
 GAA CAG TGG CTC AAG ATG AGA AGC GTA TTG AGA CAA AGA ATT CTG AAA CCG AAA GAT GTG 540  
  
 A I Y S G E V D Q V I A D L I K R I Y L 200  
 GCC ATT TAT TCT GGA GAA GTC GAC CAA GTT ATT GCT GAC TTA ATT AAA AGA ATC TAC CTC 600  
  
 L R S Q A E D G E T V T N V N D L F F K 220  
 CTC AGG AGC CAG GCA GAA GAT GGA GAA ACC GTG ACC AAT GTC AAT GAT CTT TTC TTC AAA 660  
  
 Y S M E G V A T I L Y E S R L G C L E N 240  
 TAT TCA ATG GAA GGA GTG GCC ACC ATC CTT TAT GAG AGT CGT TTG GGC TGC CTG GAA AAC 720  
  
 S I P Q L T V E Y I E A L E L M F S M F 260  
 AGC ATC CCA CAG CTG ACT GTG GAA TAC ATC GAG GCC CTG GAG CTC ATG TTT AGC ATG TTC 780  
  
 K T S M Y A G A I P R W L R P F I P K P 280  
 AAG ACC TCC ATG TAT GCA GGC GCC ATC CCC AGA TGG CTT CGC CCC TTC ATC CCA AAG CCC 840  
  
 W R E F C R S W D G L F K F S Q I H V D 300  
 TGG CGG GAA TTC TGC AGG TCC TGG GAT GGA CTC TTC AAA TTC AGC CAA ATT CAT GTT GAC 900  
  
 N K L W D I Q Y Q M D R G R R V S G G L 320  
 AAC AAG TTG TGG GAC ATA CAG TAC CAA ATG GAC CGA GGC CGG AGG GTG AGC GGG GGA CTT 960  
  
 L T Y L F L S Q A L T L Q E I Y A N V T 340  
 CTC ACA TAC CTC TTC CTT AGC CAG GCT CTG ACG CTG CAG GAG ATC TAC GCC AAC GTG ACT 1020  
  
 E M L L A G V D T T S F T L S W T V Y L 360  
 GAG ATG CTG CTG GCC GGC GTC GAC ACG ACG TCC TTC ACC TTG TCT TGG ACG GTG TAC CTC 1080

FIG. 1a

L A R H P E V Q Q T V Y R E I V K N L G 380  
CTG GCA AGG CAC CCA GAA GTG CAG CAG ACG GTG TAC CGG GAG ATT GTG AAG AAT TTA GGG 1140

E R H V P T A A D V P K V P L V R A L L 400  
GAA AGG CAT GTT CCA ACT GCA GCT GAT GTC CCC AAG GTC CCG CTG GTC AGA GCT CTC CTT 1200

K E T L R L F P V L P G N G R V T Q E D 420  
AAG GAA ACC CTG AGG CTG TTT CCA GTG CTG CCA GGG AAC GGC CGG GTC ACC CAG GAA GAC 1260

L V I G G Y L I P K G T Q L A L C H Y A 440  
CTG GTT ATT GGC GGG TAT CTG ATT CCG AAA GGC ACC CAG CTG GCC CTT TGC CAC TAT GCC 1320

T S H Q D E N F P R A K E F R P E R W L 460  
ACA TCG CAC CAG GAT GAG AAC TTC CCT CGG GCC AAG GAG TTC CGA CCT GAG CGC TGG CTG 1380

R K G D L D R V D N F G S I P F G H G V 480  
CGG AAA GGA GAC TTA GAT AGA GTT GAC AAT TTT GGA TCC ATC CCC TTT GGT CAT GGG GTT 1440

R S C I G R R I A E L E I H L V V I Q L 500  
CGC AGC TGC ATA GGG CGG AGA ATT GCA GAA CTG GAG ATT CAC CTC GTC GTG ATC CAG TTG 1500

L Q H F E I K T S S Q T N A V H A K T H 520  
CTT CAA CAT TTT GAG ATC AAA ACA TCT TCT CAG ACC AAT GCT GTT CAT GCA AAA ACC CAC 1560

G L L T P G G P I H V R F V N R K \* 538  
GGG CTC CTG ACG CCA GGG GGG CCC ATC CAC GTG CGA TTT GTT AAC AGA AAG TAA 1614

←SEQ ID NO:3↑

GCCTAGATTTTAAACCTGGGCTGATGTAGCAGACCAGCTCGCCGACACACAGTGGGTATTTGTGTTCGCTGATCACCGT

GGAGAAGGAAAGCGATGTGCGCTAAAGGCTGTCTTGTATAGACTGGCCTCCCAGGTCCTGGGACACTTGTAATCTTTA

TGCAAAGTAATGTAAAAAGGTTGCTATTTTACTGGTGCATACCAGAAGTTGCCCTTTCTTTGGGGGAAACAGCTGTTTA

AAAACAGTGGCAGTGAATTTTATGCTTCATACATTGNGCTAGACTCAATATTTAATGACTGGCAGTATCCTGTGCAT

TTACTTGTACAGGGAAATGGTGGNTTACTTACAAATTCAGTTCTTC

FIG. 1b

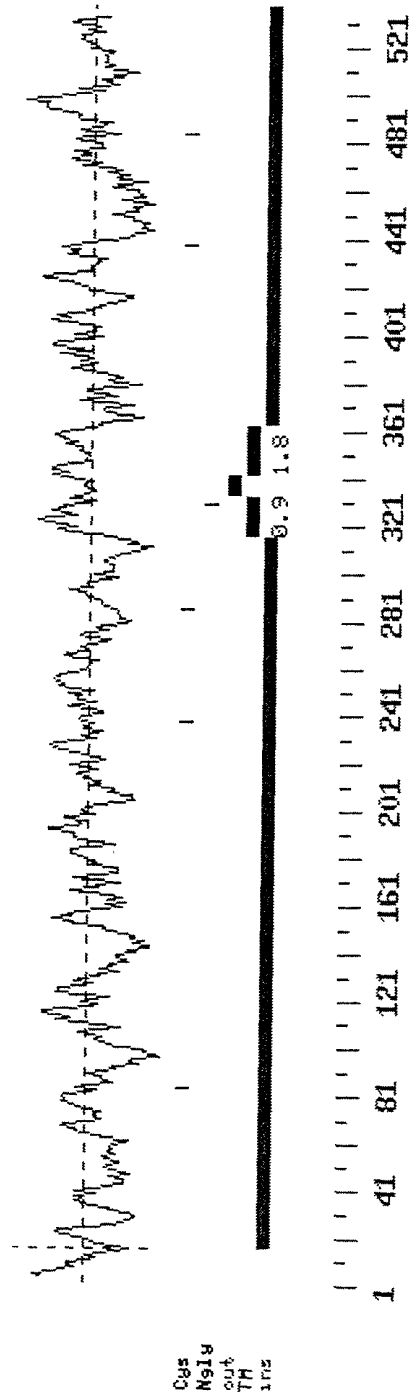


FIG. 2

```

*->PPgPpplPliGnll.qlgrappgph.sltklrkakrYGkpvftlyl SEQ ID NO:4
  PgP l ++ ++ g+ i h++ +k + + YG ++f ++
27439 69 MGPRTLANLAEFFcRDGF--SRI-HeIQQKHTR-E-YG-KIFKSHF 109

  GprpvVvltgpeavkevLidkgeeFakgRgdfnptfpwlskgyre.....
  Gp+ vV + + ++v +vL +g ++ R + + w+ yr+ +++
27439 110 GPQFVVSIA DRDMVAQVLRAEGAAPQ--RANM---ESWR--EYRDlrgra 152

  qglifsdnGpkWrklRrFslltlrfHFgmGaysKrsqkleePriqeeard
  +gl+ + +G++W k+R l +r +++ +++++ +++++ d
27439 153 TGLISA-EGEQWLKMRS--VLRQRI-LKP-----KDVAIYSGEVDQVIAD 193

  Lverlrk...eqa.gspi.DitellarlaplnvIcsllFGvrfdylr...
  L++r+ ++++++g+++++ +l++ ++ ++ ++ +l+ r++l+++
27439 194 LIKRIYLLrsQAEDGETVtNVNDLFF-KYSMEGVATILYESRLGCLensi 242

  peDp.eflklidklInemfdrvspwhqllldifP.fLlrylpgslfrkafk
  p + e++ +++ l+ mf ++s+ ++ +P++L+++ p++ +r+ +
27439 243 PQLTVEYIEALE-LMFsMF-KTSM--YAG-AIPrWLRPFIPKP-WREFC- 285

  aakd.lkdyldklieerretlepagdpRrld.i....gflDslLleakr
  + +d+l+++ + + + ++ + + D+ ++ ++g+l + l++++
27439 286 RSWDgLFKFSQIHVDNKLW--DI-QYQM--DrGrrvsgGLL-T-YLFLSQ 328

  eggnpkselsdeelaatvldllfAGteTTsstLswaLylLakhPevqakl
  l+ e+ a+v ++l AG++TTs tLsw++yLLa+hPevq+ +
27439 329 -----ALTLQEIYANVTEMLLAGVDTTSTLSWTVYLLARHPEVQQTv 371

  reEideviGrdrsptydvDaraqmPYLdAvIkEtLRlypvvPlllpRvat
  + Ei +G pt D +++ P++ A +kEtLRl+pv+P + +Rv+
27439 372 YREIVKNLGERHVPtAA-D-VPKVPLVRALLKETLRlFPVLP-GNGRVTQ 418

  kDteiPdGylIPKGtlVivnlyslhrDpkvfpnPeeFdPeRFL....de
  +D +i +GylIPKGt+ ++ y+ + ++ fp++ eF+PeR+L++++ d
27439 419 EDLVI-GGYLIPKGtQLALCHYATSHQDENFPRAKEFRPERWlrkgdldr 467

  ngkfkksyaflPFGaGpRnClGerlArmElflfLatlLqrFPelelavpp
  + + +PFG+G+R+C+G+r+A++E+ l++++lLq+F e++++ +
27439 468 ----VDNFGSIPFGHGVRSICGRRIAELIHLVVIQLLQHF-EIKTSSQT 512

  gdipsltpkpelglpskpplykvqlrpa<-*
  ++ + + ++l++ +p++ +r +
27439 513 NAVH-AKTH--GLLTPGGPIH--VR-F 533

```

FIG. 3

Query: 146 RDLRGRATGLISAEGEQWLKMRSVLRQRILKPKDVAIYSGEVDQVIADLIKRIYLLRSQA 205  
 RD RG A GL++AEGE+W ++RS L +++KP++V Y ++++V D ++R+ +R Q  
 Sbjct: 11 R DYRG EAYGLLTAEGE EWQLRSALNP KLMKPQEVKNYIPKLNEVSQDFVERLRKMRDQG 70 SEQ ID NO:5

Query: 206 E-DGETVTNVNDLFFKYSMEGVATILYESRLGCL E-NSIPQLTVEYIEALELMFSMFKTS 263  
 + GE V + + +K++ E + T+L+ RLGCL E N++ ++I+A++ SMF T+  
 Sbjct: 71 QGQ GELVEDFAEELYKWAFESICTVLFGR LGCLEENNVDPEAQKFIDAVK---SMFHTT 127

Query: 264 MYAGAI PRWL-RPFIPKPWREFCRSWDGLFKFSQIHVDNKLWDIQYQMDRG 313  
 + +P L R F K W++ R+WD +F Q ++D L ++ + G  
 Sbjct: 128 VPMNMPP ELWRYFKTKTKD HVRWDQIFDVCQKYIDEALERLEKESQSG 178

FIG. 4

Query: 71 GPRTL ANLAEFFCRDGF SRIHE-IQQKHTREY GKIFKSHFG-PQFVVS IAD RDMVAQVLR 128  
 G + L + G + +H+ I +H ++YG IF+ G Q V ++ +++ V +  
 Sbjct: 64 GLPVVGT LVDLIAAGGATHLHKYIDARH-KQYGP IFRERLGGTQDAVFVSSANLMRGVFQ 122 SEQ ID NO:6

Query: 129 AEGAAPQRANMESWREYRDLRGRATGLISAEGEQWLKMRSVLRQRILKPK---DVAIYS 184  
 EG PQ ++W Y GL EG +WL R +L + +L DV I S  
 Sbjct: 123 HEGQYPQHPLPD AWTLYNQQHACQ RGLFFMEGAEWLHNRILNRLLLNGNLNWM DVHIES 182

Query: 185 GEVDQVIADLIKRIYLLRSQ---AEDGET---VTNVNDLFFKYSMEGVATILY-ESRLG 236  
 + + D KR + AE GE + + +++S+E + I++ S L  
 Sbjct: 183 --CTRRMVDQWKRRTA EAAAIP LAESGEIRSYELPLLEQQLYRWSIEVLCCIMFGTSVLT 240

Query: 237 CLENSIPQLTVEYIEALELMFSMFKTS MYAGAI PRWLRPFIPKP-WREFCRSWDGLFKFS 295  
 C + Q +++Y +++ +F+ S P L + P WR+F + D + +  
 Sbjct: 241 CPKI---QSSLDYFT--QIVHKVFEHSSRLMTFPPRLAQILRLPIWRDFE ANVDEV LREG 295

Query: 296 QIHVDNKLWDIQYQMDRGRVSGGLLT YLFSQALTLQEIYANV-TEMLLAGVDTTSTFL 354  
 +D+ I+ Q D+ R L Y L A ++ + ++++A DTT+F+  
 Sbjct: 296 AAIDHC---IRVQEDQRRPHDEAL--YHRLQAADVP GDMIKRIFVDLVIAAGDTTAFSS 350

Query: 355 SWTVYLLARHPEVQQT VYREIVKNLGERHVPTAADVPKVPLVRALLKETLRLF 407  
 W ++ L++ P +QQ + +E N R L+ L+KE+LRL+  
 Sbjct: 351 QWALFALSKEPRLQORLAKERATN-DSR-----LMHGLIKESLRLY 390

FIG. 5

Query: 66 LAAMPGPRTL ANLAEFFCRDGF SRIHEIQQKHTREY GKIFKSHFGPQFVVS IAD RDMVAQ 125  
 LA +PGP TL L + FC+ +HE+Q YG ++ S FG Q V++A +V Q  
 Sbjct: 37 LADIPGP GTLRLFLFQ LFCCKGYLLHLHELQVLGKARYGPMWMS SFGTQRTVN LASPPLVEQ 96 SEQ ID NO:7

Query: 126 VLRAEGAAPQRANMESWREYRDLRGRA 152  
 V+R EG P R + E W+E+R RA  
 Sbjct: 97 VMRQEGKYPVRCSFEPWKEHRRRHQRA 123

FIG 6

TOP SECRET

FIG. 7

FIG. 8

FIG. 9

Query: 88 SRIHEIQKHKTREYKGIFKSHFGPQFVVSIA DRMVAQVLR AEGAAPQRANMESWREYRD 147  
 + + + + + + + E K FG + + + + + + + AQ + + E + N + Y  
 Sbjct: 52 TELYDLYIRESMEKYGAVKFFFGSRWNILVSRSEYLAQIFKDEDTFAKSGNQKKI-PYSA 110 SEQ ID  
 NO:11

Query: 148 LRGRATG--LISAEGEQWLKMRSVLRQRILKPKDVAIYSGEVDQVIADLIKRIYLLRSQA 205  
 L TG +ISA G W R+ + + D I+ + + + LIK L  
 Sbjct: 111 LAAY-TGDNVISAYGAVWRNRYNAVNTNGLQHFD DAPIFKNA--KILCTLIKNRLL----- 162

Query: 206 EDGETVTNVNDLFFKYSMEGVATILYESRLGCLENSIPQLTVEYIEALELMFS-MFKTSM 264  
 +G+T + L + + + + + + G L + I + +F F T  
 Sbjct: 163 -EGQTSIPMGPLSQRMALDNISQVALGFDFGALT HEKNAFHEHLIRIKKQIFHPFFLTFP 221

Query: 265 YAGAIPRWLRPFIPKWPREFCRSWDGLFKFSQIHVDNKLWDIQYQMDRGRRVSGGLLTYL 324  
 + +P P K + + + + L K Q + N Y+ + + + L+  
 Sbjct: 222 FLDVLP I---PSRKKAFKDVVSFREL LVKRVQDEL VNN-----YKFEQTTFAASDLIR-A 272

Query: 325 FLSQALTLQEIYANVTEMLLAGVDTTSTLSWTVYLLARHPEVQQT VYREIVKNLGERHV 384  
 + + + + + N+ +L+AG + + +YLLA++ Q R+ V + +  
 Sbjct: 273 HNNEIIDYKQLTDNIVIIILVAGHENPQLLFNSSLYLLAKYSNEWQEKLRKEVNGITD--- 329

Query: 385 PTAADV PKVPLVRALLKETLRLFPVL 410  
 P + +PL+ A L E +R++P L  
 Sbjct: 330 PKG--LADLPLLNAFLFEVVRMY PPL 353

FIG. 10

Query: 396 VRALLKETLRLFPVLPNGRVTQEDLVIGGYLIPKGTQLALCHYATSHQDENFPRAKEFR 455  
 +R + ET+RL + P R + + + + GY IP T + + +  
 Sbjct: 23 LRQVQDETIRLSTLAPWAARYSDKKVTVCGYITIPAKTPMIHALGVGLKNKTVWENTDSWD 82 SEQ ID  
 NO:12

Query: 456 PERWLRKGDLD RVDNFGSIPFG-HGVRSCIGRRIAELEIHLVVIQLLQHFEIKTSSQTNA 514  
 P+R+ G R ++F PFG H R C G + E+ + LL FEI  
 Sbjct: 83 PDRFSPNGR--RGNDF--CPFGVHSRRKCPGYLFSYFEVGVFASILLSRFEIVPVEGQTV 138

Query: 515 VHAKTHGLLT-PGGPIHVR FVNRK 537  
 + + HGL+T P I + +RK  
 Sbjct: 139 I--QVHGLVTEPKDDIKIYIRSRK 160

FIG. 11

094304064650

Query: 344 LAGVD'TSFTLSWTVYLLARHPEVQQT'VYREIVKNLGERHVP'PTAADVPKVPLVRALLKET 403  
+AG+DT + +L++ +Y + H E + E L PTA + + PL+ + ET  
Sbjct: 247 IAGMDTAANSLAFVLYRMHLHSEFLPALRAE-ADALFRDGPPTAEALGRSPLLHRFVMET 305 SEQ ID  
NO:13  
Query: 404 LRLFPVLP 411  
LR+ P+ P  
Sbjct: 306 LRVHPIAP 313

FIG. 12a

Query: 69 MPGPRTLANLAEFFCRDGF'SRIHEIQQKHTREYGKIFKSHFGPQFVVS'ADRDMAQVLR 128  
+P P L + AE G ++ + + YG +F+ Q + +A + ++  
Sbjct: 2 VPAPPFLGHAAEM----GTIKLRPFLTRCYQAYGPVFQLTVP'GQKITVLAGPEANLFAMK 57 SEQ ID  
NO:14  
Query: 129 AEGAAPQRANMESWREYRDLRGRATGLISAEGEQWLKMR'SV 169  
EG R+ +E+WR+ G +IS +G + R V  
Sbjct: 58 -EGHRVLR'S-LEAWRDNDHEMGSDRSMISLDGAEH'RAYRRV 96

FIG. 12b

Query: 153 TGLISAEGEQWLKMR'SVLRQRILKPKDVAIYSGEVDQVIADLIKRIYLLRSQAEDGE--- 209  
T IS GEQW KMR V+ I+ PK + G+ + +L+ ++ + ++E  
Sbjct: 2 T'VAISPYGEQWKMRKVITTEIMSPKRLNWLLGKRTEEADNLVAYVHNMCQKSETNNKHG 61 SEQ ID  
NO:15  
Query: 210 TVTNVNDLFFKYSMEGVATILYESR-LG----CLENSIP-QLTVEYIEALELMFSMFKTS 263  
V +V D+ Y V +++ R G ++ P E+++A+  
Sbjct: 62 AVIDVRDVVRHYCHNVMMR'FMFGRRHFGKGTZFSDDGGPGPEEKEHMDAIFTALDCLYAF 121  
Query: 264 MYAGAI'PRWLR 274  
+ IPRWLR  
Sbjct: 122 CVSDYIPR'WLR 132

FIG. 13a

Query: 458 RWLRKGDLD 466  
RWLR DLD  
Sbjct: 129 RWLRGWDLD 137 SEQ ID NO:16

FIG. 13b